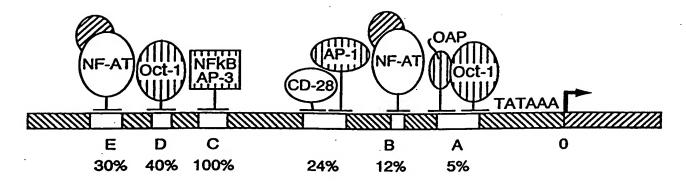
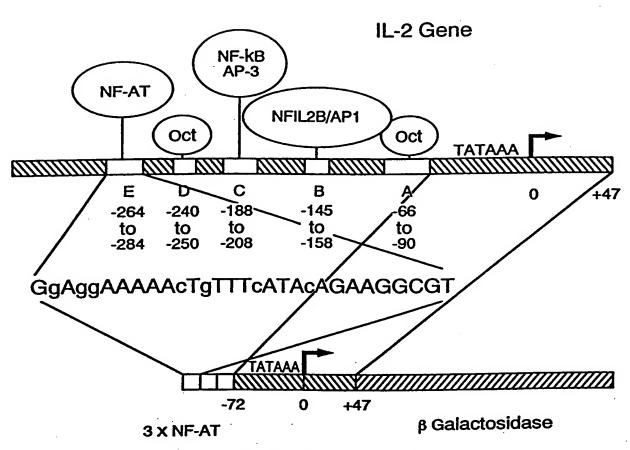


Fig. 1



IL-2 Enhancer

Fig. 2



NF-AT Gal Fusion Gene

Fig. 3

FP J+ J- K+ K- F+ F- H T F C J- J+

H





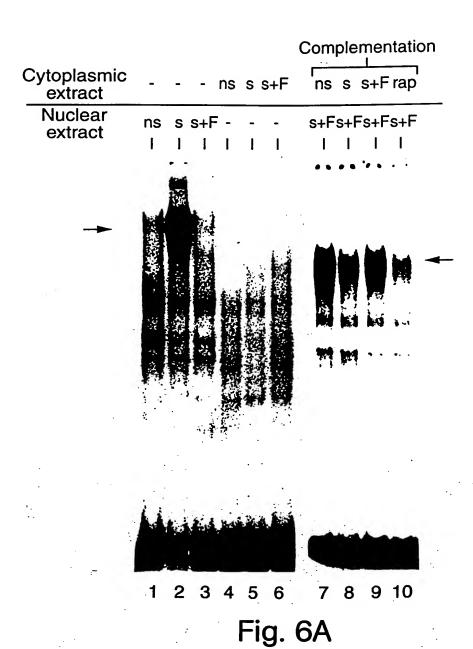
Fig. 4

Cytoplasmic extract	-	-	ns	ns	ns	ns
anisomycin	-	-	-	. -	+	+
Nuclear extract	s+F	s+F	s+F	s+F	s+F	s+F
anisomycin	-	+	-	+	-	+
	1	1	1	1	1	1



1 2 3 4 5 6

Fig. 5



Competing oligos

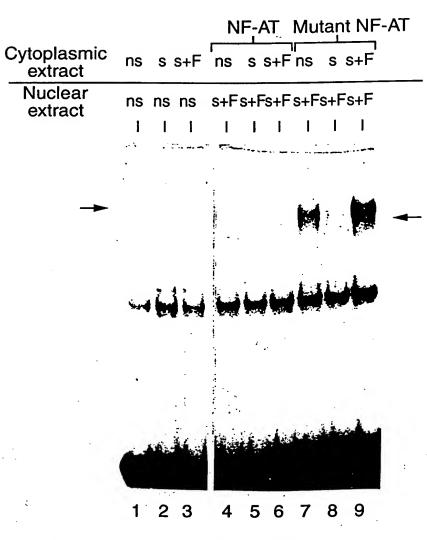


Fig. 6B

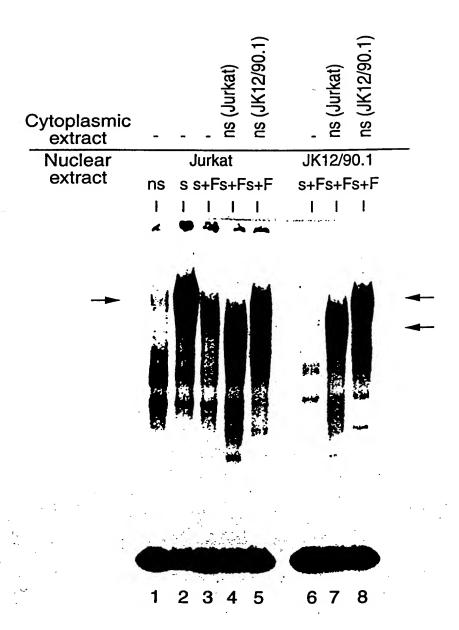
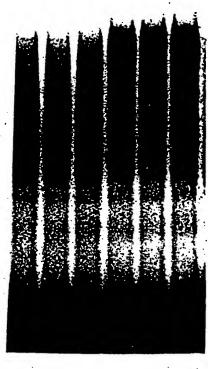


Fig. 6C

Cytoplasmic extract	ns	s	s+C	ns	S	s+C	
Nuclear extract	ns	ns	ns	s+C	s+C	s+C	-
extract	I	1	I	ı	ı	l	
	1 4	# .	. 1	#40	-		



1 2 3 4 5 6

Fig. 6D

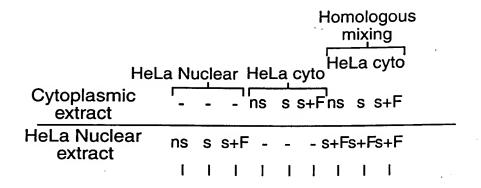
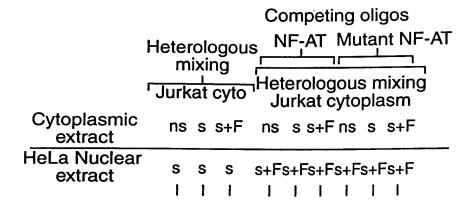
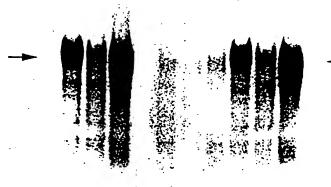




Fig. 7A







1 2 3 4 5 6 7 8 9

Fig. 7B

Heterologous mixing

Cytoplasmic extract

Jurkat Nuclear s+Fs+Fs+F extract

| 1 | 1 |



Fig. 7C

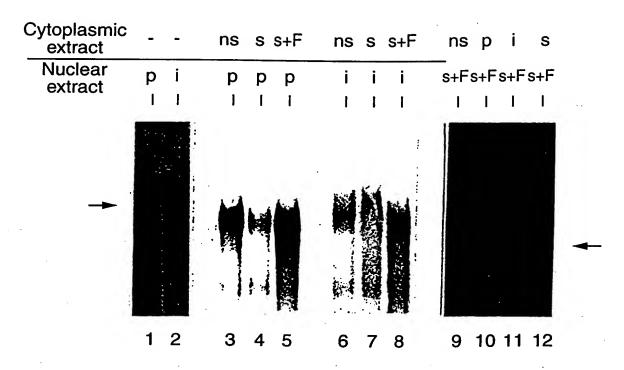


Fig. 8

Normalized fold induction	PMA/ionomycin + FK506	- nonstim 	
Test G-less →	~		~
AdMLP internal control		5 • •	

1 2 3

Fig. 9A

nonstimPMA/ionomycin+FK506

NF-AT/LacZ mRNA ─►

1 2 3

Fig. 9B

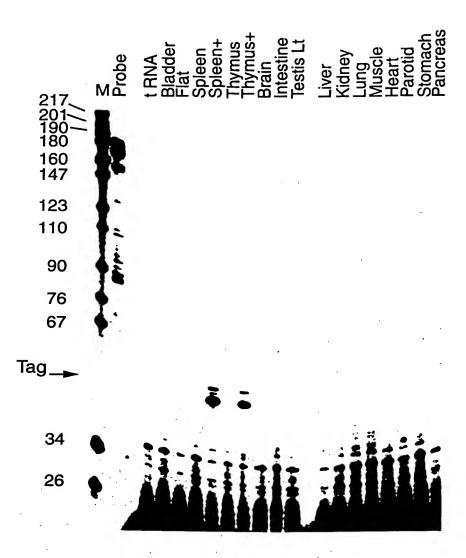


Fig. 10

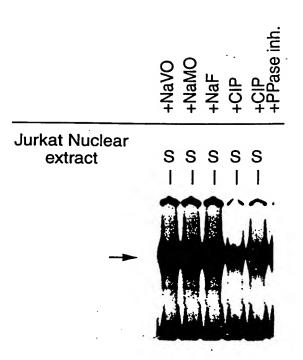




Fig. 11

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	10	30	50	70	90	
	gaatteegeagggegeggg 110	caccggggcgcgggcaggg 130	rctcggagccaccgcgcagg 150	gtcctagggccgcggccggg 170	gccccgccacgcgcgcacacgcccc 190	
	tegatgacttteeteeggg 210	egegegggegetgageeeg 230	gggcgagggctgtcttccc 250	eggagacccgaccccggcag 270	gcgcgggggggccacttctcctgtg 290	
1	cctccgcccgctgctccac	teccegeegeegegegegegegegegegegegegegegeg		ccagtcccttccaagtttc PVPSKFI 370	ccacttggccctgcggctgcggtct P L G P A A A V F 390	21
22	tcgggagaggagaacttt G R G E T L 410	ggggcccgcgcgcgcgcc G P A P R A 430	ggcggcaccatgaagtcag G G T M K S A 450	 gggaggaagaacactatgg A E E E H Y G 470	gctatgcatcctccaacgtcagccc Y A S S N V S P 490	54
55	cgccctgccgctccccacg	gcgcactccaccctgccgg	ccccgtgccacaaccttca	gacctccacaccgggcato	catecegeeggeggateaceceteg I P P A D H P S 590	87
88	•	acggtgggcccgcgggcta	cttcctctcctccggccac	accaggcctgatggggccc	ectgccctggagagtcctcgcatcg	121
122	agataacctcgtgcttggg I T S C L G 710	cctgtaccacaacaataac LYHNNN 730	cagtttttccacgatgtgg Q F F H D V E 750	aggtggaagacgtcctccc	tagetecaaacggteceetecae SSKRSPST 790	154
155	ggccacgctgagtctgccca A T L S L P S 810	agcctggaggcctacagag 5 L E A Y R D 830	acccctcgtgcctgagccc PSCLSP 850	ggccagcagcctgtcctcc A S S L S S 870	cggagctgcaactcagaggcctcc R S C N S E A S 890	187
188	tcctacgagtccaactactc S Y E S N Y S 910	egtacccgtacgcgtccccc Y P Y A S P 930	ccagacgtcgccatggcag Q T S P W Q 9	tctccctgcgtgtctccca S P C V S P K 970	agaccacggaccccgaggagggct T T D P E E G F 990	221 .
222	ttccccgcgggctgggggcc P R G L G A 1010	tgcacactgctgggttccc C T L L G S I 1030	cogcagcactccccctccac Q H S P S T 1050	cetegeeeegegeeagegte SPRASV 1070	cactgaggagagctggctgggtgc TEESWLGA	254
255	ccgctcctccagacccgcgt R S S R P A S	ccccttgcaacaagaggaa PCNKR 1130		gcagccgccctactcacccc Q P P Y S P I 1170	caccactegecacgeegteeeg H H S P T P S P 1190	287
288	cacggctccccgcgggtcag H G S P R V S	cgtgaccgacgactcgtgg V T D D S W	ttgggcaacaccacccagt L G N T T Q N	acaccagctcggccatcgt TSSAIV	tggccgccatcaacgcgctgacca A A I N A L T T	321

Fig. 12A

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	1210	1230	1250	1270	1290
32	ccgacagcagcetggacct DSSLDL 1310	.gggagatggcgtccctgtca G D G V P V R 1330	agtcccgcaagaccaccct S R K T T L 1350	ggagcagccgccctcagtgg E Q P P S V A	gcgctcaaggtggagcccgtcgg A L K V E P V G 354 1390
355	ggaggacctgggcagcccc 5 E D L G S P 1410	ccgccccggccgacttcgc P P P A D F A 1430	gcccgaagactactcctcti PEDYSS1 1450	ttccagcacatcaggaaggg F Q H I R K G 1470	gegettetgegaceagtacetg G F C D Q Y L 387 1490
388	gcggtgccgcagcacccct BAVPQEPY 1510	accagtgggcgaagcccaag Q W A K P K 1530	cccctgtcccctacgtccta P L S P T S Y 1550	acatgagcccgaccctgccc M S P T L P 1570	gccctggactggcagctgccgt A L D W Q L P S 421 1590
422	cccactcaggcccgtatga H S G P Y E 1610	gcttcggattgaggtgcagc L R I E V Q P 1630	ccaagtcccaccaccgagcc KSHHRA 1650	ccactacgagacggagggca H Y E T E G S 1670	gccggggggccgtgaaggcgtc R G A V K A S 454 1690
455	ggccggaggacaccccatc A G G H P I 1710	gtgcagctgcatggctactt V Q L H G Y L 1730	ggagaatgagccgctgatgc ENEPLMI 1750	etgcagcttttcattgggac , Q L F I G T 1770	ggeggaegaecgeetgetgege A D D R L L R 487 1790
488	ccgcacgccttctaccagg P H A F Y Q V 1810	tgcaccgcatcacagggaag HRITGK 1830	accgtgtccaccaccagcca F V S T T S H 1850	cgaggctatcctctccaac E A I L S N 1 1870	accaaagtcctggagatcccac T K V L E I P L 521 1890
522	tcctgccggagaacagcat L P E N S M 1910	gcgagccgtcattgactgtgc R A V I D C A 1930	cggaatcctgaaactcaga G I L K L R 1950	aactccgacattgaacttc N S D I E L R 1970	ggaaaggagagacggacatcgg K G E T D I G 554 1990
555	gaggaagaacacacgggtad R K N T R V F 2010	ggctggtgttccgcgttcac LVFRVH 2030	gtcccgcaacccagcggcc V P Q P S G R 2050	gcacgctgtccctgcaggtg T L S L Q V 2070	ggcctccaaccccatcgaatgc A S N P I E C 587 2090
588	tcccagcgctcagctcagga S Q R S A Q B 2110	gctgcctctggtggagaagc LPLVEKQ 2130	agagcacggacagctatcc S T D S Y P 2150	ggtcgtgggcgggaagaaga V V G G K K M 2170	atggtcctgtctggccacaact N V L S G H N F 621 2190
622	tcctgcaggactccaaggtc L Q D S K V 2210	attttcgtggagaaagcccc I F V E K A P 2230	agatggccaccatgtctgg D G H H V W I 2250	yagatggaagcgaaaactga B M E A K T D 2270	ccgggacctgtgcaagccgaa R D L C K P N 654 2290
655	ttctctggtggttgagatcc S L V V E I P 2310	cgccatttcggaatcagagg PFRNQR 2330	ataaccagccccgttcacgt I T S P V H V 2350	cagtttctacgtctgcaac S F Y V C N 2370	gggaagagaagcgaagccag G K R K R S Q 687 2390
688	taccagcgtttcacctacct YQRFTYL	tcccgccaacggtaacgccal P	cctttctaaccgtaagccgt F L T V S R	gaacatgagcgcgtggggt E H E R V G C	gctttttctaaagacgcagaa PP 7716

Fig. 12B

20/31

2470

2490

acgacgtcgccgtaaagcagcgtggcgtgttgcacatttaactgtgtgatgtcccgttagtgagaccgagccatcgatgccctgaaaaggaaaaggaaaag 2510 2530 2550 2570 2590

cagccaaggggaaaacatggctcttctgctccaaaaaactgagggggtcctggtgtgcatttgcaccctaaagctgcttacggtgaaaaggcaaataggt 2710 2730 2750

Fig. 12C

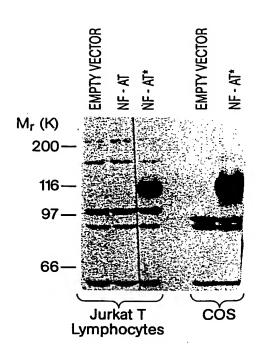
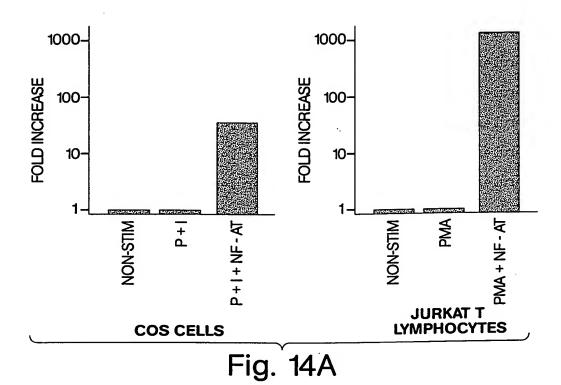


Fig. 13



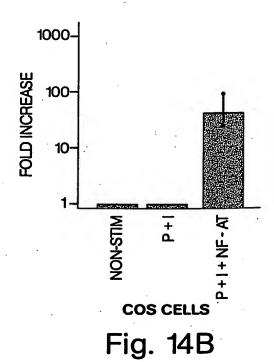


Fig. 15A

542

DMDORSAL C-REL NFKB p50 NFKB p65 NFATC NFATC	LNSVRLCFQVFMESEQK LNVVRLCFQVFL-PDEH LSVVRLMFTAFL-PDST LNAVRLCFQVTV-RDPA NTRVRLVFRVHV-PQPS
DMDORSAL	VSE
c-REL NFKB p50	SNPIYDNRAPNTAELRICRV-NKNCGSVRGGD SDAIYDSKAPNASNLKIVRM-DRTAGCVTGGE
	LSHPIFDNRAPNTAELKICRV-NRNSGSCLGGDEIFLLC
.	SNPI-ECSQRSAHELPMVE
DMDORSAL C-REL	KVAKEDISVRFFEERNGQ-SVWEAFGDFQHTDVHKQTAITFKTPRYHTLD KVQKDDIEVRFVLNDWEAKGIFSQADVHRQVAIVFKTPPYCK-A
NFKB p50	FGDFSPTDVHRQFAIVFKTPKYK
NFKB p65	KVQKEDIEVYFTGPGWEARGSFSQADVHRQVAIVFRTPPYADPS
NFATC	HNFLQDSKVIFVEKAPDGHHVWEMEAKT-DRDLCKPNSLVVEIPPFRNQR
NFATP	QNFTAESKVVFMEKTTDGQQIWEMEATV-DKDKSQPNMLFVEIPEVRNKH
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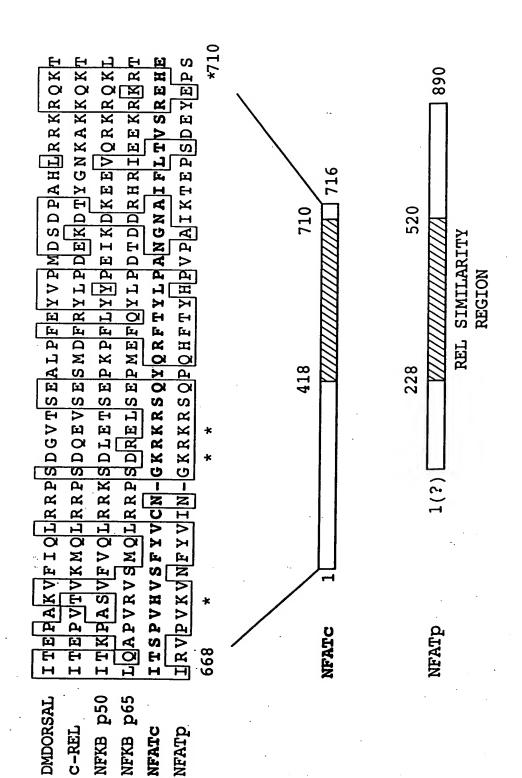


Fig. 15C

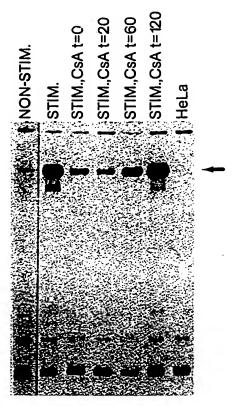


Fig. 16A

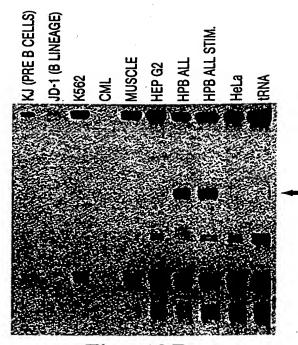


Fig. 16B

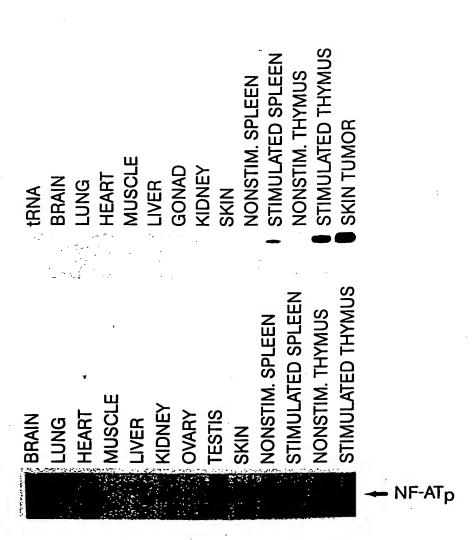
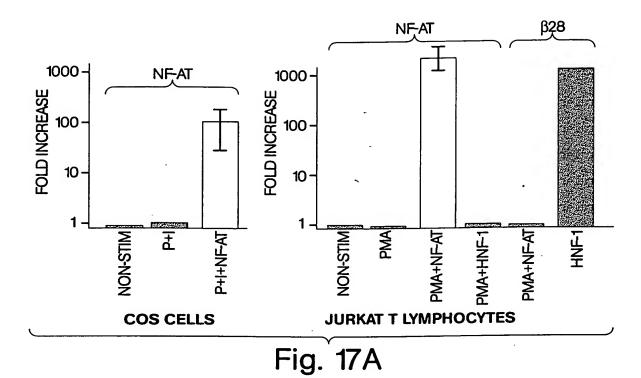
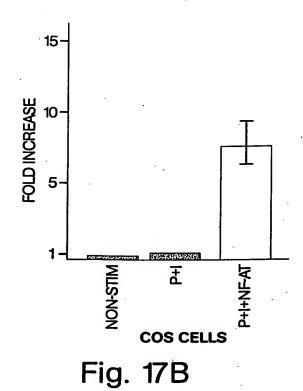
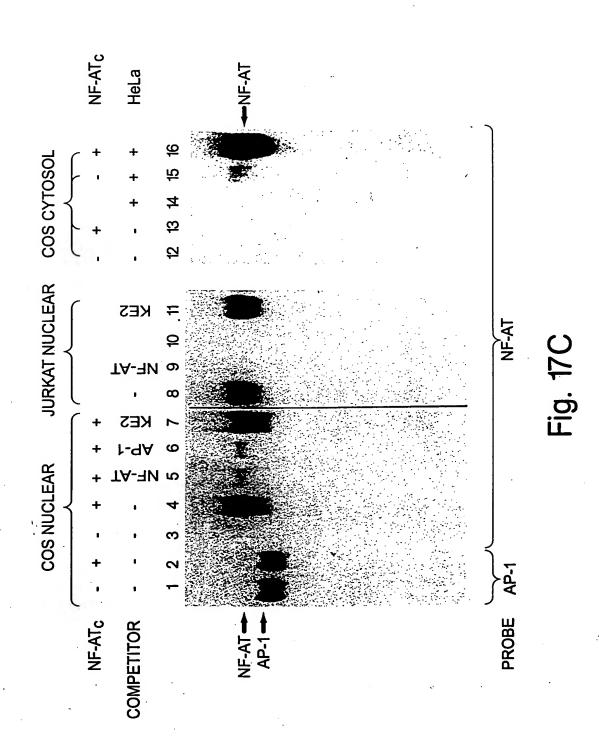


Fig. 16C







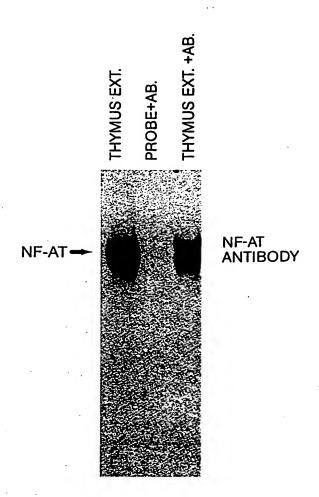


Fig. 17D

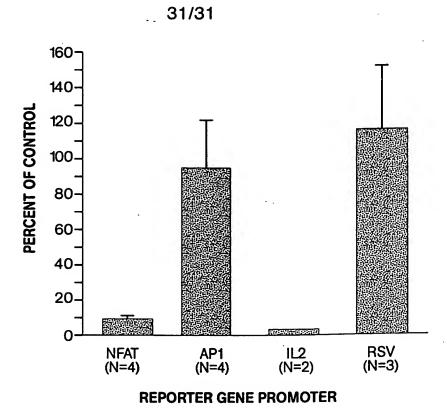


Fig. 18

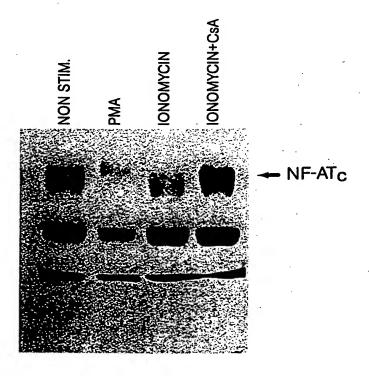


Fig. 19